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Perfect score:

Run on:

Scoring table: Sequence:

Searched:

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Sequence 22, Application US/08924759
Patent No. 596229
GENERAL INFORMATION
TITLE OF INVENTION: PLANT GLUTATHIONE-S-TRANSFERASE
TITLE OF INVENTION: PRIVES
CORRESPONDENCE ADDRESS:
ADDRESSEE: B.I. DU PONT DE NEMOURS AND COMPANY
STREET: 1007 MARKET STREET
CITY: WILLINGTON
STATE: DELAWARE
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.50 INCH
COMPUTER: IBM PC COMPATIBLE
COMPATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMPUTER: TELECOMPUTER: CASACTER FILE
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US-09-248-335-64

US-09-248-335-66

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US-09-248-335-66

US-09-248-335-52

US-09-248-335-52

US-09-248-335-52

US-09-248-335-52

US-09-248-335-52

US-09-247-3738-26

US-09-247-3738-26

US-09-247-3738-36

US-09-247-3738-36

US-09-247-3738-36

US-09-247-3738-36

US-09-247-3738-36

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TOPOLOGY: not relevant
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CLONE: CEB5.PK0049.A11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: protei
ORIGINAL SOURCE:
TISSUE TYPE: MAIZE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: not
RESULT 1
US-08-924-759-22
389.5
383.5
383.5
383.5
373.5
372.5
373.5
373.5
365.5
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360.5
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Sequence 22, 1
Sequence 12, 1
Sequence 12, 1
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48,
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/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/pcTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/pcTUS_COMB.pep:*
          GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-248-747-6
US-09-246-715-6
US-09-247-747-20
US-09-247-747-20
US-09-247-715-20
US-09-247-715-20
US-09-247-712-12
US-09-247-747-12
US-09-247-747-18
US-09-247-747-18
US-09-296-715-12
US-09-247-747-18
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US-08-525-507-2
US-09-248-335-48
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Listing first 45 summaries
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Maximum DB seq length: 200000000
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Database

3

Gaps

4;

73.1%; Score 857; DB 2; Length 224; 72.9%; Pred. No. 9.7e-91; Live 27; Mismatches 30; Indels

Conservative

Similarity

Query Match Best Local Simil Matches 164; (

Sequence Sequence Sequence Sequence

-09-248-335-40 -09-248-335-54 -09-248-335-74 US-09-248-335-60 US-08-924-747-24

6601.5 6011.5 6011.5 6011.5 6611.5 643.2 643.2 643.0 6

857 818.5 818.5 681.5 681.5 664.5 664.5 664.5 664.5

Result Ş. Sequence Sequence

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                                                                                                                               KIKGEPQAQARAEMIDILKTIDGALGDKPFFGGDKFGFVDAAFAPFTAWFHSYERYGEFS 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  59 VLLHDGRAVNESLIILQYLEEAFPD-APALLPSDPYARAQARFWADYVDKKVYDCGSRLW 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KIKGEPQAQARAEMLDILKTLDGALGDKPFFGGDKFGFVDAAFAPFTAWFHSYERYGEFS 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 120 KLKGDGQAQARAEMVEILRTLEGALGDGPFFGGDALGFVDVALVPFTSWFLAYDRFGGVS 179
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                 MAGEK--GLVLLDFWVSPFGQRVRIALAEKGLPYEYAEEDLMAGKSDRLLRANPVHKKIP
                                                              59 VLLHDGRAVNESLIILQYLEEAFPD-APALLPSDPYARAQARFWADYVDKKVYDCGSRLW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 MAGEK -- GLVLLDFWVSPFGQRVRIALAEKGLPYEYAEEDLMAGKSDRLLRANPVHKKIP
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APPLICANT: O'KEEF, DANIEL

TITLE OF INVENTION: PLANT GLUTATHIONE-S-TRANSFERASE ENZYMES

TITLE OF INVENTION: PLANT GLUTATHIONE-S-TRANSFERASE ENZYMES

FILE REFERENCE: CL-1128-A

CURRENT APPLICATION NUMBER: US/09/248,335

CURRENT APPLICATION NUMBER: 08/924,759

EARLIER FILING DATE: 1997-September-05

NUMBER OF SEQ ID NOS: 74

SEQ ID NO 22

LENGTH: 224

ENGLEN 224
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 12, Application US/08924759
Patent No. 596229
GENERAL INFORMATION:
APPLICANT: MCGONIGLE, BRIAN
APPLICANT: O'KEEFE, DANIEL
TITLE OF INVENTION: PLANT GLUTATHIONE-S-TRANSFERASE
TITLE OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: E.I. DU PONT DE NEMOURS AND COMPANY
STREET: 1007 MARKET STREET
                                                                                                                                                                                                                                                                                                                Sequence 22, Application US/09248335 Patent No. 6096504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity ....
Matches 164; Conservative
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                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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ORGANISM: maize
                                                                                                                                                                                                                                                                                               US-09-248-335-22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                           COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.50 INCH
COMPUTER: 1BM PC COMPATIBLE
OPERATING SYSTEM: MICROSOFT WORD FOR WINDOWS 95
SOFTWARE: MICROSOFT WORD VERSION 7.0A
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/924,759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      69.8%; Score 818.5; DB 2
69.5%; Pred. No. 2.6e-86;
tive 31; Mismatches 33
DELAWARE : UNITED STATES OF AMERICA
                                                                                                                                                                                                                                     CL-1128
                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: FLOYD, ILNDA ARAMETHY
REGISTRATION NUMBER: 33,692
REFERENCE/DOCKET NUMBER: CL-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 302-892-8112
TELEPHONE: 302-773-0164
                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS: not relevant
TOPOLOGY: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: protein
ORIGINAL SOURCE:
TISSUE TYPE: maize
TIMEDIATE SOURCE:
CLONE: ceb5.pk0051.f8
US-08-924-759-12
                                                                                                                                                                                                                                                                               TELEFAX: 302-773-0164 INFORMATION FOR SEQ ID NO: 12:
                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 225 amino acid
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 69.5%
Matches 157; Conservative
                                                                                                                                                       FILING DATE:
CLASSIFICATION: 435
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67 VNESLIILQYLEEAFPDAPALLPSDPYARAQARFWADYVDKKVYDCGSRLWKLKGEPQAQ 126
                                                  67 VNESLIILQYLEEAFPDAPALLPSDPYARAQARFWADYVDKKVYDCGSRLWKLKGEPQAQ 126
                                                                        127 araemidiiktidgalgdkpffggdkfgfydaafapftawfhsyerygefslpeyapkia 186
                                                                                                                                              7 LVLLDFWVSPFGQRVRIALAEKGLPYEYAEEDIMAGKSDRLLRANPVHKKIPVLLHDGRA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COPERATION SYSTEM: MICROSOFT WORD FOR WINDOWS 95
SOFTWARE: MICROSOFT WORD VERSION 7.0A
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/296,715
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: FLOYD, LINDA AXAMETHY
REGISTRATION NUMBER: 33,692
REFERENCE/POCKET UNBER: 33,692
REFERENCE/POCKET UNBER: 31,692
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1NFORMATION:
TELECOMMUNICATION 1NFORMATION:
TELECOMMUNICATION 10164
                                                                                                                                                                                                                     187 AWAKRCGERESVAKSLYSPDKVYDFIGLLKKKYGIE 222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NEMOURS AND COMPANY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 683.5; DB 4
Pred. No. 8.9e~71;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: DELAWARE
COUNTRY: UNITED STATES OF AMERICA
COUNTRY: BEADABLE FORM:
MEDIUM TYPE: DISKETTE, 3.50 INCH
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
                                                                                                                                                                                                                                                                                                                        Sequence 6, Application US/09296715
Patent No. 6171839
                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: MCGONIGLE, BRIAN
APPLICANT: O'KEFE, DANIEL
TITLE OF INVENTION: SOYBEAN GI
TITLE OF INVENTION: ENZYMES
NUMBER OF SEQUENCES: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: E.I. DU PONT DE
STREET: 1007 MARKET STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             not relevant
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amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: protein
ORIGINAL SOURCE:
TISSUE TYPE: SOYBEAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 1007 MARK
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Best Local Similarity
Matches 130; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino a STRANDEDNESS:
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MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                         US-09-296-715-6
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                                                                                                                                                                                 PVLLHDGRAVNESLIILQYLEEAFPDA-PALLPSDPYARAQARFWADYVDKKVYDCGSRL 116
                                                                                                                                                                                                                                WKLKGEPQAQARAEMLDILKTLDGALGDKPFFGGDKFGFVDAAFAPFTAWFHSYERYGEF 176
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                                                                                                      1 MAGE--KGLVLLDFWVSPFGQRVRIALAEKGLPYEYAEEDLMAG-KSDRLLRANPVHKKI
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                                                    Gaps
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              Length 225;
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                                                                                                                                                                                                                                                                                                       SLPEVAPKIAAWAKRCGERESVAKSLYSPDKVYDFIGLLKKKYGIE 222
                                                                                                                                                                                                                                                                                                                           33; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: MGONIGLE, BRIAN
APPLICANT: O'KEEFE, DANKEL
TITLE OF INVENTION: SOYBEAN GLUTATHIONE-S-TRANSFERASE
TITLE OF INVENTION: BNZYMES
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSE: E.I. DU PONT DE NEMOURS AND COMPANY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: MICROSOFT WORD FOR WINDOWS 95 SOFTWARE: MICROSOFT WORD VERSION 7.0A CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/924,747
              DB 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             E: E.I. DU PONT DE NEMOURS AND COMPANY 1007 MARKET STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  58.3%; Score 683.5; DB 3 60.2%; Pred. No. 8.9e-71; Live 27; Mismatches 58
                             ; Pred. No. 2.6e-86; 31; Mismatches 33
              Score 818.5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: FLOXD. LINDA AXAMETHY
REGISTRATION NUMBER: 33,692
REFERENCE/DOCKET NUMBER: CL-1108
TELECOMUNICATION INFORMATION:
TELEPHONE: 302-812
TELEPHAX: 302-773-0164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 UNITED STATES OF AMERICA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.50 INCH
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: MICROSOFT WORD
                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 6, Application US/08924747
Patent No. 6063570
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: protein
ORIGINAL SOURCE:
              69.88;
                               69.58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 219 amino acids
TYPE: amino acid
STRANDEDNESS: not releva
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                  Matches 157; Conservative
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STATE: WILMINGTON
STATE: DELAWARE
COUNTRY: ITWE
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                               Best Local Similarity
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Best Local Sim
Matches 130;
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MICROSOFT WORD VERSION 7.0A
               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/924,747
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                                                                           CLASSIFICATION: 435
ATTORNEY AGENT INFORMATION:
NAME: FLOYD. LINDA AXAMETHY
RECISTRATION NUMBER: 33,692
REFERENCE/DOCKET NUMBER: CL-11
TELECHONE: 302-892-8112
TELEPHONE: 302-892-8112
TELEFAX: 302-773-0164
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                             LENGTH: 219 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: protein
ORIGINAL SOURCE:
TISSUE TYPE: SOYBEAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Microsoft Office 97
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CLONE: SS1.PK0005.E6
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SEQ ID NO 20
LENGTH: 219
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                                      127 ARAEMIDIIKTIDGALGDKPFFGGDKFGFVDAAFAPFTAWFHSYERYGEFSLPEVAPKIA 186
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7 LVLLDFWYSPFGQRVRIALAEKGLPYEYAEEDLMAGKSDRLLRANPVHKKIPVLLHDGRA
                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: MCGONIGLE, BRIAN
APPLICANT: O'KEEFE, DANIEL
TILE OF INVENTION: SOUPEAN GLUTATHIONE-S-TRANSFERASE ENZYMES
FILE REFERENCE: CL-1108-A
CURRENT APPLICATION NUMBER: US/09/247,373B
CURRENT FILING DATE: 1999-02-10
PRIOR APPLICATION NUMBER: 08/924,747
PRIOR APPLICATION NUMBER: 08/924,747
PRIOR FILING DATE: 1997-09-05
NUMBER OF SEQ ID NOS: 56
SOFTWARE: Microsoft Office 97
LENGTH: 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       58.1%; Score 681; DB 4; Length 218; 60.6%; Pred. No. 1.7e-70; Live 27; Mismatches 56; Indels
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Patent No. 6603570

GENERAL INFORMATION:
APPLICANT: O'KERFE, DAINIEL
TITLE OF INVENTION: SOYBEAN GLUTATHIONE-S-TRANSFERASE
TITLE OF INVENTION: BOXYBEAN GLUTATHIONE-S-TRANSFERASE
TITLE OF INVENTION: BOXYBEAS
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSE: E.I. DU PONT DE NEMOURS AND COMPANY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: DISKETTE, 3.50 INCH
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: MICROSOFT WORD FOR WINDOWS 95
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                                                                                                                           184 AWAKRCLQKESVAKSLPDQQKVYEFIMDLRKKLGIE 219
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STREET: 1007 MARKET STREET
CITY: WILMINGTON
                                                                                                  187 AWAKRCGERESVAKSLYSPDKVYDFIGLLKKKYGIE 222
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                                                                                                                                                                                                                                                    Sequence 6, Application US/09247373B Patent No. 6168954
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Best Local Similarity
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ORGANISM: SOYBEAN
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                                                                                                                                                                                                                                                                                                                  127 ARAEMLDILKTLDGALGDKPFFGGDKFGFVDAAFAPFTAWFHSYERYGEFSLPEVAPKIA 186
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APPLICANT: O'KBERE, DANIEL
TITLE OF INVENTION: SOYBEAN GLUTATHIONE-S-TRANSFERASE ENZYMES
FILE REFERENCE: C1-1108-A
CURRENT APPLICATION NUMBER: US/09/247,373B
PRIOR APPLICATION NUMBER: 08/99-02-10
PRIOR APPLICATION NUMBER: 08/924,747
PRIOR FILING DATE: 1997-09-05
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                                                            Indels
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          DB 3;
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                                   .4e-68;
Query Match 56.6%; Score 664.5; Best Local Similarity 57.9%; Pred. No. 1.4e Matches 125; Conservative 31; Mismatches
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ARAEMLDILKTLDGALGDKPFFGGDKFGFVDAAFAPFTAWFHSYERYGEFSLPEVAPKIA 186
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                                                                                                                                                                                          Sequence 12, Application US/08924747
Patent No. 6063570
GENERAL INFORMATION:
APPLICANT: MCGONIGLE, BRIAN
APPLICANT: O'REEFE, DANIEL
TITLE OF INVENTION: SOYBEAN GLUTATHIONE-S-TRANSFERASE
TITLE OF INVENTION: ENZYMES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: MICROSOFT WORD FOR WINDOWS 95
SOFTWARE: MICROSOFT WORD VERSION 7.0A
CURRENT APPLICATION DATA:
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                                                                          AWAKRCGERESVAKSLYSPDKVYDFIGLLKKKYGIE 222
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STREET: 1007 MARKET STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 639; DB 3;
Pred. No. 1.2e-65;
                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                      UNITED STATES OF AMERICA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/924,747
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MEDIUM TYPE: DISKETTE, 3.50 INCH
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY AGENT INFORMATION:
NAME: FLOYD, LINDA AXAMETHY
REGISTRATION NUMBER: 33,692
REFRENCE/DOCKET NUMBER: CL-1:
TELECOMMUNICATION INFORMATION:
TELEFAM: 302-773-0164
INFORMATION FOR SEQ ID NO: 12:
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not relevant
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Best Local Similarity 55.9%
Matches 124; Conservative
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                                                                                                                                                                                                                                                                           TITLE OF INVENTION: EN UTILE OF INVENTION: EN NUMBER OF SEQUENCES: 3 CORRESPONDENCE ADDRESS:
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CLONE: SE6.PK0048.D7
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CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                    WILMINGTON
                                                                                                                                                                                                                                                                                                                                                                                                  STATE: DELAWARE
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US-08-924-747-12
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                                                                                                                      VNESLIILQYLEEAFPDAPALLPSDPYARAQARFWADYVDKKVYDCGSRLWKLKGEPQAQ 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     VNESLIILQYLEEAFPDAPALLPSDPYARAQARFWADYVDKKVYDCGSRLWKLKGEPQAQ 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5 VVLLDFWPSPFGMRVRIALAEKGIEYEYKEEDL-RNKSPLLLQMNPVHKKIPVLIHNGKP 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7 LVLLDFWVSPFGQRVRIALAEKGLPYEYAEEDLMAGKSDRLLRANPVHKKIPVLLHDGRA 66
                                                      ARAEMLDILKTLDGALGDKPFFGGDKFGFVDAAFAPFTAWFHSYERYGEFSLPEVAPKIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: MGGONTGLE, BRIAN
APPLICANT: MGGONTGLE, DANIEL
TITLE OF INVENTION: SOYBEAN GLUTATHIONE-S-TRANSFERASE
TITLE OF INVENTION: RAZYMES
NUMBER OF INVENTION: BAZYMES
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPATIENT IN PC COMPATIBLE
COMPATING SYSTEM: MICROSOFT WORD FOR WINDOWS 95
SOFTWARE: MICROSOFT WORD VERSION 7.0A
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/296,715
                                                                                                                                                                                          187 AWAKRCGERESVAKSLYSPDKVYDFIGLLKKKYGIE 222
                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: E.I. DU PONT DE NEMOURS AND COMPANY
STREET: 1007 MARKET STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 56.6%; Score 664.5; DB 4; Best Local Similarity 57.9%; Pred. No. 1.4e-68; Matches 125; Conservative 31; Mismatches 59;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   UNITED STATES OF AMERICA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DISKETTE, 3.50 INCH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CL-1108
                                                                                                                                                                                                                                                                                        Sequence 20, Application US/09296715 Patent No. 6171839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: FLOYD, LINDA AXAMETHY
REGISTRATION NUMBER: 33,692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: CI
TELECOMMUNICATION INFORMATION:
TELEPHONE: 302-892-8112
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not relevant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 302-773-0164
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: protein ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOYBEAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SS1.PK0005.E6
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STATE: DELAWARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amino acid
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CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19898
                                                                                                                                                                                                                                                        RESULT 10
US-09-296-715-20
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                                 29
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121 GEPQAQARAEMLDIIKTILDGALGDKPFFGGDKFGFVDAAFAPFTAWFHSYERYGEFSLPE 180
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 MAGEKGLVLLDFWVSPFGQRVRIALAEKGLPYEYAEEDLMAGKSDRLLRANPVHKKIPVL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 216;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      181 VAPKIAAWAKRCGERESVAKSLYSPDKVYDFIGLLKKKYGIE 222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: MCGONICLE, BRIAN
APPLICANT: O'KEEFE, DANIEL
TITLE OF INVENTION: SOYBEAN GLUTATHIONE-S-TRANSFERASE
TITLE OF INVENTION: BNZYMES
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 19988
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.50 INCH
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: MICROSOFT WORD FOR WINDOWS 95
SOFTWARE: MICROSOFT WORD VERSION 7.0A
APPLICATION NUMBER: US/08/924,747
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: E.I. DU PONT DE NEMOURS AND COMPANY
STREET: 1007 MARKET STREET
CITY: WILLINGTON
STATE: DELAWARE
COUNTRY: UNITED STATES OF AMERICA
                                                                                                                                                                                                                                                                                                                                                                                                   54.5%; Score 639; DB 4;
55.9%; Pred. No. 1.2e-65;
tive 33; Mismatches 59
                         CL-1108
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                    REFERENCE/DOCKET NUMBER: CL-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 302-8912
TELEFAX: 302-773-0164
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 216 amino acids
                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: protein
ORIGINAL SOURCE:
TISSUE TYPE: SOYBEAN
REGISTRATION NUMBER: 33,692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: FLOYD, LINDA AXAMETHY
REGISTRATION NUMBER: 33,692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: CI
TELECOMMUNICATION INFORMATION:
TELEPHONE: 302-892-8112
TELEFAX: 302-773-0164
                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 55.9%
Matches 124; Conservative
                                                                                                                                                                                                                                                                              ; IMMEDIATE SOURCE:
; CLONE: SE6.PK0048.D7
US-09-296-715-12
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US-08-924-747-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LHDGRAVNESLIILQYLEEAFPDAPALLPSDPYARAQARFWADYVDKKVYDCGSRLWKLK 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            58 IHNGKPICESLVAVQYIEEVWHDRNPLLPSDPYQRAQARFWADFVDNKIFDLGRKIWTSK 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 MAGEKGLVILIDFWVSPFGQRVRIALAEKGLPYEYAEEDLMAGKSDRLLRANPVHKKIPVL 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: WCGONIGLE, BRIAN
APPLICANT: O'KERE, DANIEL
TITLE OF INVENTION: SOYBEAN GLUTATHIONE-S-TRANSFERASE ENZYMES
FILE REFERENCE: CL-1108-A
CURRENT PIPLICATION NUMBER: US/09/247,373B
CURRENT FILING DATE: 1999-02-10
PRIOR APPLICATION NUMBER: 08/924,747
PRIOR FILING DATE: 1997-09-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          54.5%; Score 639; DB 4; Length 216; 55.9%; Pred. No. 1.2e-65; tive 33; Mismatches 59; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 12, Application US/09296715
Patent No. 6171839
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: MCGONICLE, BRIAN
APPLICANT: O'KEEFE, DANIEL
TITLE OF INVENTION: SOYBEAN GLUTATHIONE-S-TRANSFERASE
TITLE OF INVENTION: ENZYMES
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
ADDRESSEE: E.I. DU PONT DE NEMOURS AND COMPANY
STREET: 1007 MARKET STREET
CITY: WILMINGTON
STATE: DELAWARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             175 ECPKFVAWAKRCLQKDSVAKSLPDQHKVYEFIMDIRKKFDIE 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 181 VAPKIAAWAKRCGERESVAKSLYSPDKVYDFIGLLKKKYGIE 222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 1988
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.50 INCH
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: MICROSOFT WORD FOR WINDOWS 95
SOFTWARE: MICROSOFT WORD VERSION 7.0A
APPLICATION NUMBER: US/09/296,715
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                UNITED STATES OF AMERICA
                                                           RESULT 12
US-09-247-373B-12
: Sequence 12, Application US/09247373B
: Patent No. 6168954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: FLOYD, LINDA AXAMETHY
                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 56
SOFTWARE: Microsoft Office 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches 124; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: UNTERESTRE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; ORGANISM: SOYBEAN US-09-247-373B-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Mic
SEQ ID NO 12
LENGTH: 216
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US-09-296-715-12
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TYPE: PRT

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66 AVNESLIILQYLEEAFPDAPALLPSDPYARAQARFWADYVDKKVYDCGSRLWKLKGEPQA 125
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               66 AVNESLIILQYLEEAFPDAPALLPSDPYARAQARFWADYVDKKVYDCGSRLWKLKGEPQA 125
                                                                                                                                                                                                                                                                                                                                                                                                            126 QARAEMLDILKTLDGALGDKPFFGGDKFGFVDAAFAPFTAWFHSYERYGEFSLPEVAPKI 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6 GLVLLDFWVSPFGQRVRIALAEKGLPYEYAEEDLMAGKSDRLLRANPVHKKIPVLLHDGR 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  6 GLVLLDFWVSPFGQRVRIALAEKGLPYEYAEEDLMAGKSDRLLRANPVHKKIPVLLHDGR 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       51.3%; Score 601.5; DB 4; Length 200; 55.3%; Pred. No. 2.2e-61; Live 32; Mismatches 55; Indels 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: MCGONIGLE, BRIAN
APPLICANT: O'KEEFE, DANIEL
TITLE OF INVENTION: SOCREAN GLUTATHIONE-S-TRANSFERASE ENZYMES
FILE REFERENCE: CL-1108-A
CURRENT APPLICATION NUMBER: US/09/247,373B
CURRENT FILING DATE: 1999-02-10
PRIOR PPLICATION NUMBER: 08/924,747
RIOR FILING DATE: 1997-09-05
NUMBER OF SEQ ID MOS: 56
SOFTWARE: Microsoft Office 97
                                                                                                                                                                                                                                DB 3; Length 200;
                                                                                                                                                                                                                                                                     55; Indels
                                                                                                                                                                                                                              Query Match 51.3%; Score 601.5; DB 3
Best Local Similarity 55.3%; Pred. No. 2.2e-61;
Matches 109; Conservative 32; Mismatches 55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-247-373B-18
Sequence 18, Application US/09247373B
Patent No. 6168954
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        VAWAKRCMOREAVSKSL 199
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183 VAWAKRCMQREAVSKSL 199
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAWAKRCGERESVAKSL 202
                                                         SS: not relevant
not relevant
SEQUENCE CHARACTERISTICS:
LENGTH: 200 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 55.38
Matches 109; Conservative
                                                                                           MOLECULE TYPE: protein ORIGINAL SOURCE: TISSUE TYPE: SOYBEAN
                                                                                                                                                  ; IMMEDIATE SOURCE:
; CLONE: SS1.PK0002.F7
US-08-924-747-18
                                     TYPE: amino ac
STRANDEDNESS:
TOPOLOGY: not
MOLECULE TYPE: I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: SOYBEAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-247-373B-18
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Search completed: July 1, 2003, 09:00:58 Job time: 16 secs

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GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on:

July 1, 2003, 08:53:35; Search time 21 Seconds (without alignments) 1159.124 Million cell updates/sec

US-09-508-710-2

1173 1 MAGEKGLVLLDEWVSPFGGR.....rspdkvydfigllkkkygie 222 Perfect score: Sequence:

BLOSUM62 Scoring table:

424699 seqs, 109646833 residues Gapop 10.0 , Gapext 0.5 Searched:

424699

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Total number of hits satisfying chosen parameters:

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database:

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/cgn2_6/ptodata/1/pubpaa/USO9_NEW_PUB.pep:*
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/cgn2_6/ptodata/1/pubpaa/USO9_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/USO9_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/USO9_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/USO9_NEW_PUB.pep:*
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//cgn2_6/ptodata/1/pubpaa/USO0_NEW_PUB.pep:*
//cgn2_6/ptodata/1/pubpaa/USO0_NEW_PUB.pep:* Published_Applications_AA:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

		df				
Result No.	Score	Query Match	Query Match Length DB	DB	a	Description
	405.5	34.6	236	10	US-09-765-213A-9	fund 0 equenos
2	362.5	30.0		-	115-00-765-0138-8	
6	334		•	9 6	HG-00-765-2138-2	
**	331.5	28.3	-	2	TS-09-765-2138-7	Sequence 2, Appli
ĸ	320.5	27.3	224	10	US-09-765-213A-11	-
φ	319	27.2		10	US-09-765-213A-4	4
7	309	26.3		10	US-09-765-213A-6	Sequence 6, Appli
æ	291	24.8		10	US-09-765-213A-10	20,
6	235	20.0		10	US-09-050-010-7	7. A
10	198	16.9		10	US-09-741-669-351	Sequence 351, App
11	172.5	14.7		6	US-09-854-133-194	
12	172.5	14.7		6	US-10-225-273-4	Sequence 4, Appli
13	172.5	14.7		10	US-09-738-973-194	Sequence 194, App
14	172.5	14.7		10	US-09-925-302-653	
15	167.5	14.3		5	US-09-854-133-204	
16	167.5	14.3		10	US-09-738-973-204	Sequence 204, App
17	129.5	11.0		10	0S-09-020-010-6	Sequence 6, Appli
. 18	124	10.6		5	US-09-798-889-164	Sequence 164, App
19	121.5	10.4		6	US-09-769-787-68	•

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23, A 4105, A 4114, 4614, 3951, 3951, 3951, 3958, 403, 355, 403, 403, 403, 403, 403, 403, 403, 403	Sequence 55, Appl Sequence 180, App
9 US-09-964-899-23 9 US-10-106-698-414 10 US-09-925-297-761 10 US-09-731-872-403 10 US-09-731-872-403 10 US-09-731-872-403 9 US-110-102-806-54 9 US-110-102-806-54 10 US-09-925-310-928 10 US-09-925-310-928 10 US-09-925-310-928 10 US-09-925-310 10 US-09-926-692-1310 10 US-09-796-692-1310 10 US-09-796-692-2311 10 US-10-040-862-1311 10 US-10-040-862-1311 10 US-10-040-862-1311 10 US-09-925-300-1649 10 US-10-040-862-2051 10 US-09-925-300-1649 10 US-10-040-862-2311 10 US-09-925-300-1649 10 US-09-925-300-1649 10 US-10-072-033-13	10 US-09-833-745-55 9 US-09-981-876-180
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ALIGNMENTS

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Sequence 9, Application US/09765213A

Before W. US20020079846A1

GENERAL INFORMATION:
APPLICANT: Racchini, Peter J

ITLE OF INVENTION: NO. US20020079846A1e1 Glutathione-S-Transferase Nucleic Acids

TILE OF INVENTION: Polypeptides and Methods of Use

FILE REFERENCE: 22542-001

CURRENT FILING DATE: 2001-01-17

PRIOR APPLICATION NUMBER: 60/176708

PRIOR PILING DATE: 2000-01-18

PRIOR FILING DATE: 2000-01-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 34.6%; Score 405.5; DB 10; Best Local Similarity 42.3%; Pred. No. 7.5e-35; Matches 96; Conservative 40; Mismatches 80;
                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 9
LENGTH: 236
                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Picea mariana
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US-09-765-213A-9
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Sequence 7, Application US/09765213A

Fatent No. US20020079846A1

GENERAL INFORMATION:

APPLICANT: Facchini, Peter J

APPLICANT: Facchini, Peter J

TITLE OF INVENTION:

FILIE REFERENCE: 22542-001

CURRENT APPLICATION NUMBER: US/09/765,213A

CURRENT FILING DATE: 2001-01-17

PRIOR FILING DATE: 2000-01-18

NUMBER OF SEQ ID NOS: 11

SOFTWARE: Patentin Ver. 2.1
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Patent No. US20020079846A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Facchini, Peter J
TILLE OF INVENTION: No. US20020079846A1el Glutathione-S-Transferase Nucleic Acids
TITLE OF INVENTION: Polypeptides and Methods of Use
TILLE REFERENCE: 22542-001
CURRENT APPLICATION NUMBER: US/09/765,213A
CURRENT FILING DATE: 2001-01-17
PRIOR FILING DATE: 2000-01-18
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn Ver: 2.1
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                                                                                   123 DAEERKAAIEQAIAAFGILEEAYQKISKG----KDFFGGEKIGYVDIAFGCYYGWIRVTE 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    116 VGAKDDEGKMAAVGKLAMECLAILEETFQKSSKGLG---FFGGETIGYLDIACSALLGPIS 172
       63 HGDKPICESMIIVQYIDDVWASAGHSIIPSDPYDASIARFWATYIDDKFFPSLMGIAKSK 122
                                                     121 GEPQAQARAEM----LDIL-----KTLDGALGDKPFFGGDKFGFVDAAFAPFTAWFHSYE 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         60 LLHDGRAVNESLIILQYLEEAFPDAPALLPSDPYARAQARFWADYVDKKVYDCGSRLWKL 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 120 KG----EPQAQARAEMLDILKTLD-----GALGDKPFFGGDKFGFVDAAFAPFTAWFH 168
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         169 SYERYG--EFSLPEVAPKIAAWAKRCGERESVAKSLYSPDKVYDFIGLLKKKYGIE 222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                  172 RYGEFSL--PEVAPKIAAWAKRCGERESVAKSLYSPDKVYDFIGLLKKKYG 220
                                                                                                                                                                                              179 KMNGIKLFDEEKVPGLIKWAEKFCADETVKSVMPETDALMEF --- AKKIFG 226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 10;
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Best Local Similarity 34.7%; Pred. No. 4.7e-27;
Matches 82; Conservative 46; Mismatchee σ
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US-09-765-213A-11
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LENGTH: 225
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                                                                                     GENERAL INFORMATION:
APPLICANT: Facchini, Peter J
TITLE OF INVENTION: NO. US20020079846Alel Glutathione-S-Transferase Nucleic Acids and
TITLE OF INVENTION: Polypeptides and Methods of Use
TITLE OF INVENTION: Polypeptides and Methods of Use
FILE REFERENCE: 22442-001
CURRENT APPLICATION NUMBER: US/09/765,213A
CURRENT APPLICATION NUMBER: 60/176708
PRIOR PILING DATE: 2000-01-18
NUMBER OF SEQ ID NOS: 11
SEQ ID NO 8
LENGTH: 229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2, Application US/09765213A
Patent No. US20020079846A1
General No. US20020079846A1
General Information:
APPLICANT: Facchini, Peter J
TILLE OF INVENTION: No. US20020079846A1e1 Glutathione-S-Transferase Nucleic Acids and
TILLE OF INVENTION: Polypeptides and Methods of Use
FILE REFERENCE: 22542-001
CURRENT APPLICATION NUMBER: US/09/765,213A
CURRENT FILING DATE: 2001-01-17
PRIOR PAPLICATION NUMBER: 60/176708
PRIOR FILING DATE: 2000-01-18
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       62 HNGAPVCESMIILQYIDEVFASTGPSLLPADPYERAIARFWVAVDDKLV--APWRQW-LR 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              121 G---EPQAQARAEMLDILKTLDGALGD----KPFFGGDKFGFVDAAFAPFTAWFHSYERY 173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 30.9%; Score 362.5; DB 10; Best Local Similarity 42.3%; Pred. No. 2.5e-30; Matches 93; Conservative 33; Mismatches 79;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28.5%; Score 334; DB 10; 35.1%; Pred. No. 2.7e-27; tive 42; Mismatches 88;
                                              Sequence 8, Application US/09765213A Patent No. US20020079846A1
                                                                                                                                                                                                                                                                                                                                                                                                                          ; ORGANISM: Aegilops squarrosa
US-09-765-213A-8
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Best Local Similarity 35.18
RESULT 2
US-09-765-213A-8
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LENGTH: 233
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APPLICANT: Facchini, Peter J
TITLE OF INVENTION: No. US20020079846Alel Glutathione-S-Transferase Nucleic Acids
TITLE OF INVENTION: Polypeptides and Methods of Use
FILE REFERENCE: 22542-001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               63 HGDKPICESMIIVQYIDDVWASAGHSIIPSDPYDASIARFWATYIDDKFFPSLMGIAKSK 122
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 123 DABEKKAAIEQAIAAFGILEBAYQKTSKG----KDFFGXEKIGYXDIAFGCYXGWIRVTE 178
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CCATION: (157)...(157)

OTHER INFORMATION: Wherein Xaa is any amino acid as defined in the OTHER INFORMATION: Specification

NAME/KET: VARIANT

LOCATION: (163)...(163)

OTHER INFORMATION: Wherein Xaa is any amino acid as defined in the OTHER INFORMATION: specification

NAME/KET: VARIANT

LOCATION: (171)...(171)

OTHER INFORMATION: Wherein Xaa is any amino acid as defined in the OTHER INFORMATION: Wherein Xaa is any amino acid as defined in the OTHER INFORMATION: Wherein Xaa is any amino acid as defined in the
                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: VARIANT
LOCATION: (58)..(58)
OTHER INFORMATION: Wherein Xaa is any amino acid as defined in the OTHER INFORMATION: Specification
NAME/KEY: VARIANT
LOCATION: (61)..(61)
OTHER INFORMATION: Wherein Xaa is any amino acid as defined in the OTHER INFORMATION: Specification
NAME/KEY: VARIANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION: (189)...(189)...OTHER INFORMATION: Wherein Xaa is any amino acid as defined in the OTHER INFORMATION: Specification OTHER INFORMATION: Description of Artificial Sequence: Consensus OTHER INFORMATION: Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          93; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      41; Mismatches
CURRENT APPLICATION NUMBER: US/09/765,213A CURRENT FILING DATE: 2001-01-17
PRIOR PAPLICATION NUMBER: 60/176708
PRIOR FILING DATE: 2000-01-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/765,213A CURRENT FILING DATE: 2001-01-17 PRIOR APPLICATION NUMBER: 60/176708 PRIOR FILING DATE: 2000-01-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: specification NAME/KEY: VARIANT
                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 26.3%;
Best Local Similarity 33.3%;
Matches 77; Conservative
                                                                                                                                               NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-765-213A-10
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APPLICANT: Facchini, Peter J
APPLICANT: Facchini, Peter J
TITLE OF INVENTION: No. US20020079846Alel Glutathione-S-Transferase Nucleic Acids and
TITLE OF INVENTION: Polypeptides and Methods of Use
TITLE OF INVENTION: Polypeptides and Methods of Use
FILE REFERENCE: 22342-001
CURRENT APPLICATION NUMBER: US/09/765,213A
CURRENT PILING DATE: 2001-01-17
PRIOR APPLICATION NUMBER: 60/176708
PRIOR FILING DATE: 2000-01-18
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.1
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Patent No. US20020079846A1
GENERAL INFORMATION:
APPLICANT: Facchini, Peter J
TITLE OF INVENTION: No. US20020079846A1e1 Glutathione-S-Transferase Nucleic Acids and
FILE REPERENCE: 25542-001
                                                                                                             7;
                                                                                                                                                                                                                                                                                                                    62 HDGRAVNESLIILQYLEEAFPDAPALLPSDPYARAQARFWADYVDKKVYDCGSR-LWKLK 120
                                                                                                                                                                                                                                                                                                                                                      121 GEPQAQARAEMLDILKTLDGALGDKPFFGGDKFGF-VDAAFAPFTAWFHSYERYGEFSL- 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    121 EKEREKNVEETYEALOFLENELKDKKFFGGEEFGLDIAAVFIAF - WIPIFQEIAGLOLF 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HDGRAVNESLIILQYLEEAFPDA-PALLPSDPYARAQARFWADYVDKKVYDCGSRLWKLK 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 121 GEPQAQARAEM----LDIL-----KTLDGALGDKPFFGGDKFGFVDAAFAPFTAWFHSYE 171
                                                                                                                                                                 2 AGEKGLVLLDFWVSPFGQRVRIALAEKGLPYEYAEEDLMAGKSDRLLRANPVHRKIPVLL 61
                                                                                                                                                                                                               2 AGEKGLVLLDFWVSPFGQRVRIALAEKGLPYEYAEEDLMAGKSDRLLRANPVHKKIPVLL 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       :|:::||||||||||||::||::||::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::|:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::||:::
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                                     Length 224;
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; Pred. No. 1e-25;
43; Mismatches 90; Indels 2
                          ; Score 320.5; DB 10; Length
; Pred. No. 6.8e-26;
49; Mismatches 84; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               179 -PEVAPKIAAWAKRCGERESVAKSLYSPDKVYDFIGLLKKKY 219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            | | : | : | : | : | 179 TSEKFPILYKWSQEFINHPFVHEVLPPRDPLFAY---FKARY 217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 4, Application US/09765213A Patent No. US20020079846A1
                          27.3%;
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33.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 33.8%
Matches 78; Conservative
                                                                Best Local Similarity 35.1
Matches 78; Conservative
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; MOLECULE TYPE: protein US-09-050-010-7
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US-09-741-669-351
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US-09-854-133-194
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                                                                                                                                                                                                                                                                                             73 ILQYLEEAFPDA-----PALLPSDPYARAQARFWADYVDKKVYDCGSRLWKLKGEP--Q 124
                                                                                                                                                                                                                                                                                                                     132 AQAAEDTRAALSLLEEAFKDRSNGRAFFSGGDAAPGLLDLALGCFLPALRACERLHGLSL 191
                                                                                                                                                                                                                       14 VSPFGQRVRIALAEKGLPYEYAEEDLMAGKSDRLLRANPVHKKIPV-LLHDGRAVNESLI 72
                                                                                                                                                                                                                                             Query Match 24.8%; Score 291; DB 10; Length 240; Best Local Similarity 36.8%; Pred. No. 9.9e-23; Matches 81; Conservative 31; Mismatches 90; Indels 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 104 East Hume Avenue
CITY: Alexandria
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Irie, Ryotaro
APPLICANT: Irie, Ryotaro
APPLICANT: Hiraoka, Susumu
APPLICANT: Kasahara, Naoko
TITLE OF INVENTION: BETHOD FOR COMPARISON OF DNA BASE
NUMBER OF SEQUENCES: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                               179 PEVA--PKIAAWAKRCGERESVAKSLYSPDKVYDFIGLLK 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SCFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/050,010
FILING DATE: 30-MAR.1998
CLASSIFICATION: 435
PROOF APPLICATION NUMBER: UP 09-079586
FILING DATE: 31-MAR.1997
APPLICATION NUMBER: JP 09-079586
FILING DATE: 31-MAR.1997
ATORNEY/AGENT INFORMATION:
NAME: Beall, T., Thomas E.
REGISTRATION NUMBER: 22,410
REFERENCE/DOCKET NUMBER: ASA-707
TELECOMMUNICATION INFORMATION:
TELEFRAK: 703-684-1167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 7, Application US/09050010
; Patent No. US/20010010903A1
; GENERAL INFORMATION:
; APPLICANT: Nagai, Kelichi
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INFORMATION FOR SEQ ID NO: 7:
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.1
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM:
SOFTWARE: Patentli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS:
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Virginia
: USA
                                                                                         ; ORGANISM: Zea mays US-09-765-213A-10
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                                  SEQ ID NO 10
LENGTH: 240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-050-010-7
                                                                       TYPE: PRT
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75 QYLEEAFPDAPALLPSDPYARAQARFWADYVDKKVYDCGSRLWKLKGEPQAQARAEMLDI 134
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21 YSHQVRIVLAEKGVSFEIEHVEKD---NPPQDLIDLNP-NQSVPTLVDRELTLWESRIIM 76
                                                                                                                 1 MAGE-KGLVLLDFWVSPFGQRVRIALAEKGLPYEYAEEDLMAGKSDRLLRANPVHKKIPV 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17 FGQRVRIALAEKGLPY--EYAEEDIMAGKSDRLLRANPVHKKIPVLLHDGRAVNESLIIL 74
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Publication No. US20020183499A1
GENERAL INFORMATION:
APPLICANT: Lodes, Michael J.
APPLICANT: Mohamath, Raodoh
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: THE THERAPY AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.475C10
             20.0%; Score 235; DB 10; Length 83; 59.5%; Pred. No. 1.9e-17; tive 13; Mismatches 19; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              86; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: POSSYTH, R. Allyn
APPLICANT: Oblsen, Kari L.
APPLICANT: Obsen, Kari L.
APPLICANT: Zyskind, Judith W.
TITLE OF INVENTION: Genes identified as required for TITLE OF INVENTION: proliferation of E. coli
FILE REFERENCE: BLITRA.009A
CURRENT APPLICATION NUMBER: US/09/741,669
PRIOR APPLICATION NUMBER: US 60/173005
PRIOR FILING DATE: 1999-12-23
NUMBER OF SEQ ID NOS: 481
SOFTWARE: FASTSEQ for Windows Version 4.0
ILENGTH: 212
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30.9%; Pred. No. 5.8e-13;
Live 36; Mismatches 86;
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                                                                                                                                                                                                                   60 LLHDGRAVNESLIILQYLEEAFPD 83
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Best Local Similarity 30.99
Matches 60; Conservative
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; ORGANISM: Escherichia coli
US-09-741-669-351
                                                                  Conservative
Query Match
Best Local Similarity
**+^hes 50; Conserv
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us-09-508-710-2.rapb

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207 KEDPTVSALLTSEKDWOGFLEL 228
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                                                                                                                                                                                                                                                                                              16 PFGQRVRIALAEKGLPYEYAEEDLMAGKSDRLLRANPVHKKIPVLLH-DGRAVNESLIIL
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14.7%; Score 172.5; DB 9; Length
Best Local Similarity 28.7%; Pred. No. 3.4e-10;
Matches 58; Conservative 28; Mismatches 107; Indels
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TITLE OF INVENTION: DIARYLSULFONYLUREA BINDING PROTEINS
FILE REFERENCE: PC98608
CURRENT APPLICATION NUMBER: US/10/225,273
CURRENT FILING DATE: 2002-08-21
PRIOR FPLING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 8
SOFTWARE: PALENTIN VERSION 3.1
CURRENT APPLICATION NUMBER: US/09/854,133
CURRENT FILING DATE: 2001-05-11
NUMBER OF SEQ ID NOS: 735
SEQ TWARE: FastSEQ for Windows Version 3.0
LENGTH: 241
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Publication No. US20030027217A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Pfizer Inc.
APPLICANT: Gabel, Christopher A.
APPLICANT: Dombroski, Mark A.
APPLICANT: Geoghegan, Kieran
APPLICANT: Griffiths, Richard J.
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                                                                                                                           TYPE: PRT
CORGANISM: Homo sapien
US-09-854-133-194
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LENGTH: 241
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91 EYLDEAYP-GKKLLPDDPYEKACQKMILELFSKVPSLVGSFIRSQNKEDYAGLKEEFRKE 149
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Patent No. US20020044941A1
GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVERTION: Nucleic Acids, Proteins and Antibodies
FILE REFERENCE: PA104
CURRENT APPLICATION NUMBER: US/09/925,302
CURRENT FILING DATE: 2001-08-10
PRIOR RAPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 2000-03-08
PRIOR FILING DATE: 2009-03-08
PRIOR FILING DATE: 2009-03-08
PRIOR FILING DATE: 2009-03-08
NUMBER OF SEQ ID NOS: 896
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Mannior, Jane
APPLICANT: Mannior, Jane
APPLICANT: Kalos, Michael D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: THE THERAPY AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.47569
CURRENT APPLICATION NUMBER: US/09/738,973
CURRENT FILING DATE: 2000-12-14
NUMBER OF SEQ ID NOS: 587
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 194
TENNAME. 3.1
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                       Sequence 194, Application US/09738973 Patent No. US20020110563A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                           Algate, Paul A.
Secrist, Heather
Indirias, Carol Yoseph
Benson, Darin R.
                                                                                                     Reed, Steven G.
Henderson, Robert
Lodes, Michael J.
                                                                                                                                                                                      Fling, Steven P. Mohamath, Raodoh
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Best Local Similarity 28.7%
Matches 58; Conservative
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US-09-738-973-194
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LENGTH: 286
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APPLICANT:
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75 QYLEEAFPDAPALLPSDPYARAQARFWADYVDKKVYDCGSRLWKLKGEPQAQARAEMLDI 134
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                                                                                                                               9; Gaps
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APPLICANT: Mohamath, Raodoh
APPLICANT: Mohamath, Raodoh
APPLICANT: Henderson, Robert A.
APPLICANT: Henderson, Robert A.
APPLICANT: Henderson, Barin R.
APPLICANT: Secrist, Heather
TILLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TILLE OF INVENTION: COMPOSITIONS AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.475c10
CURRENT APPLICATION NUMBER: US/09/854.133
CURRENT FILING DATE: 2001-05-11
NUMBER OF SEQ ID NOS: 735
SOFTWARRE: PASISEQ for Windows Version 3.0
SEQ ID NO 204
LENGTH: 241
                                                                                   Query Match 14.7%; Score 172.5; DB 10; Length 286; Best Local Similarity 28.7%; Pred. No. 4.4e-10; Matches 58; Conservative 28; Mismatches 107; Indels 9;
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                    ORGANISM: Homo saptens
US-09-925-302-653
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US-09-854-133-204
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Matches 60; Conserva
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US-09-854-133-204
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TYPE: PRT
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